STATUS OF THE CLAIMS

- 1. (currently amended) [[A]] <u>An isolated</u> nucleic acid molecule encoding a protein having an amino acid sequence given in selected from the group consisting of SEQ ID NO. 2 and sequences at least 90% homologous to SEQ ID NO:2, wherein said protein <u>has</u>, or a protein with α-1,6-mannosyltransferase activity having an amino acid at least 90% homologous to the amino acid sequence of SEQ ID NO. 2, wherein said nucleic acid is derived from *Hansenula polymorpha*.
- 2. (currently amended) The <u>isolated</u> nucleic acid molecule according to claim 1, wherein the nucleic acid is designated as SEQ ID NO. 1.
- 3. (currently amended) [[A]] <u>An isolated protein which is coded by the nucleic acid of chain claim 1.</u>
- 4. (original) A recombinant vector comprising a nucleic acid molecule designated as SEQ ID NO. 1, deposited under accession number KCTC 10583BP.
- 5. (original) A *Hansenula polymorpha* Hpoch2Δ mutant strain deposited under accession number KCTC 10584BP.
- 6. (original) The *Hansenula polymorpha* Hpoch 2Δ mutant strain according to claim 5, comprising an expression vector for a sugar chain-modifying enzyme.
- 7. (currently amended) The *Hansenula polymorpha* Hpoch2Δ mutant strain according to claim 6, wherein the sugar chain-modifying enzyme is selected from the group consisting of α-1,2-mannosidase, mannosidase IA, mannosidase IB, mannosidase IC, mannosidase II, N-acetyl glucosaminyltransferase I[[,]] and N-acetyl glucosaminyltransferase and fucosyltransferase.
- 8. (original) A process for producing a recombinant glycoprotein with reduced

glycosylation using the *Hansenula polymorpha* Hpoch2 Δ mutant strain according to claim 5 <u>lacking of α -1,6-mannosyltransferase activity, wherein the recombinant</u> glycoprotein lacks further sugar-chain synthesis of Man₈ on N-linked glycosylation.

- 9. (original) The process according to claim 8, wherein the *Hansenula polymorpha* Hpoch2 Δ mutant strain comprises an expression vector for a sugar chain-modifying enzyme, wherein said sugar chain-modifying enzyme is α -1,2-mannosidase.
- 10. (canceled)
- 11. (original) A glycoprotein produced by the process of claim 8 or 9.
- 12. (new) The Hansenula polymorpha Hpoch 2Δ mutant strain according to claim 6, wherein the sugar chain-modifying enzyme is α -1,2-mannosidase.